
Sequence Listing was accepted.

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Reviewer: markspencer

Timestamp: Mon Jul 23 15:28:13 EDT 2007

Validated By CRFValidator v 1.0.2

Application No: 10531475 Version No: 1.0

Input Set:

Output Set:

Started: 2007-07-23 11:51:54.640 **Finished:** 2007-07-23 11:53:18.911

Elapsed: 0 hr(s) 1 min(s) 24 sec(s) 271 ms

Total Warnings: 21
Total Errors: 0

No. of SeqIDs Defined: 2776

Actual SeqID Count: 2776

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Input Set:

Output Set:

Started: 2007-07-23 11:51:54.640 **Finished:** 2007-07-23 11:53:18.911

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Total Warnings: 21
Total Errors: 0

No. of SeqIDs Defined: 2776

Actual SeqID Count: 2776

Error code Error Description

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Gly Tyr Phe Ser Gly Pro Trp Ile Ile Arg Ile Thr Phe Ile Leu Phe 50 60

Ala Ile Trp Trp Ala Val Gly Glu Ile Phe Arg Leu Ser Leu Leu Arg 65 70 75 80

Arg His Arg Arg Leu Leu Ser Gly Leu Asp Leu Arg Trp Gln Glu Asn 85 90 95

Val Cys Lys Trp Tyr Ile Val Ser Asn Leu Gly Phe Ala Glu Pro Cys 100 105 110

Leu Phe Leu Thr Leu Met Phe Leu Leu Arg Ala Pro Leu Lys Met Glu 115 120 125

Ser Gly Ala Leu Ser Gly Lys Trp Asn Arg Asp Thr Ala Gly Tyr Ile 130 135 140

Ser Glu Ser Arg Leu Asn Gly Gly Ser Gly Ser Tyr Val Lys Leu Pro 165 170 175

His Asp Phe Thr Arg Thr Tyr Ser Arg Val Ile Ile Asp His Asp Glu 180 185 190 Val Ala Leu Cys Thr Tyr Pro Leu Leu Ser Thr Ile Leu Leu Gly Val 195 200 205

Phe Ala Ala Val Leu Thr Ala Tyr Leu Phe Trp Leu Gly Arg Gln Ile 215 220 210

Leu Lys Leu Val Ile Asn Lys Arg Leu Gln Lys Arg Val Tyr Thr Leu 225 230 235 240

Ile Phe Ser Val Ser Ser Phe Leu Pro Leu Arg Ile Val Met Leu Cys 245 250

Leu Ser Val Leu Thr Ala Ala Asp Lys Ile Ile Phe Glu Ala Leu Ser 260 265 270

Phe Leu Ala Phe Leu Ser Leu Phe Cys Phe Cys Val Val Ser Ile Cys 275 280 285

Leu Leu Val Tyr Phe Pro Val Ser Asp Ser Met Ala Leu Arg Gly Leu 290 295 300

Arg Asp Thr Asp Asp Glu Asp Thr Ala Val Thr Glu Glu Arg Ser Gly 305 310 315 320

Ala Leu Leu Leu Ala Pro Asn Ser Ser Gln Thr Asp Glu Gly Leu Ser 325 330 335

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Asp Gly Asp Gly Lys Arg Gly Gly 65 70

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<213> Oryza sativa

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Gly Glu Gly Cys Trp Arg Ser Leu Pro Lys Ala Ala Gly Leu Leu Arg 35 40 45

Cys Gly Lys Ser Cys Arg Leu Arg Trp Met Asn Tyr Leu Arg Pro Asp 50 55 60

Leu Lys Arg Gly Asn Phe Thr Asp Asp Glu Asp Glu Leu Ile Ile Arg 65 70 75 80

Leu His Ser Leu Leu Gly Asn Lys Trp Ser Leu Ile Ala Gly Gln Leu 85 90 95

Pro Gly Arg Thr Asp Asn Glu Ile Lys Asn Tyr Trp Asn Thr His Ile 100 105 110

Lys Arg Lys Leu Leu Ala Arg Gly Ile Asp Pro Gln Thr His Arg Pro 115 120 125

Leu Leu Ser Gly Gly Asp Gly Ile Ala Ala Ser Asn Lys Arg His His 130 135 140

Arg Arg Ile Pro Tyr Pro Ser Arg Arg Arg Arg Pro Arg Arg 145 150 155 160

Ser Ser Pro Cys Glu Ala Ala Ala Ala Ala Ala Pro Gly Arg Leu Leu $165 \hspace{1.5cm} 170 \hspace{1.5cm} 175$

Gly Arg Arg Leu Pro Gln Gln Gln Arg His Asn Glu His Gly Gly Ala 180 185 190

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Val Val His Tyr Leu Cys Arg Lys Val Ala Arg Gln Pro Leu Pro Val 35 40 45

Pro Ile Ile Ala Glu Val Asp Leu Tyr Lys Leu Asp Pro Trp Asp Leu 50 55 60

Pro Glu Lys Ala Leu Phe Gly Arg Lys Glu Trp Tyr Phe Phe Thr Pro 65 70 75 80

Arg Asp Arg Lys Tyr Pro Asn Gly Ser Arg Pro Asn Arg Ala Ala Gly 85 90 95

Arg Gly Tyr Trp Lys Ala Thr Gly Ala Asp Lys Pro Val Ala Pro Lys
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Gly Ser Ala Arg Thr Val Gly Ile Lys Lys Ala Leu Val Phe Tyr Ser 115 120 125

Gly Lys Ala Pro Arg Gly Val Lys Thr Asp Trp Ile Met His Glu Tyr 130 135 140

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Ala	Ala 210	Ala	Asp	Thr	Met	Ser 215	Asp	Ser	Phe	Gln	Thr 220	His	Asp	Ser	Asp
Ile 225	Asp	Asn	Ala	Ser	Ala 230	Gly	Leu	Arg	His	Gly 235	Gly	Суз	Gly	Gly	Gly 240
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Arg Phe Ser Glu Glu Gln Ile Lys Ser Leu Glu Ser Met Phe Ala Thr 35 40 45

Gln Thr Lys Leu Glu Pro Arg Gln Lys Leu Gln Leu Ala Arg Glu Leu 50 55 60